

Pawel Rosikiewicz

Data Analyst/Analytics Translator with hands-on experience in, Molecular Biology, Industrial R&D and Product development

Data Science Machine Learning **EPFI**



Biostatistics Pipeline Dev.



R&D **Product Dev.**

Project Management





ACHIEVMENTS

Founded SwissAI, one of the lagers AI meetups in Europe www.SwissAI.org

Promoted Reproducible Data Science & Personalized Medicine

- ⇒ EPFL- Applied Machine Learning Days; https://swissai.org/amld-workshop/
- ⇒ Meetup @Silicon Valley; https://youtu.be/W624gdkDqRQ?t=1259

Interviewed 150+ High-tech Professionals.

SELECTED PROJECTS & LEADERSHIP

Product Owner & Lead Data Scientist @ SkinDiagnosticA

I developed open-source platform for analysis, and classification of medical images, with example of skin cancer. Managed, a team of 3 developers, and provided courses and workshops for users. https://simpleai.ch/skin-diagnostic-ai/

Project Manager @ *Dronistics*

8/2020 - 6/2021

9/2020 - 6/2021

I managed the project on adapting medical delivery drones to requirements from hospitals & rescue services, with the team working @Kings College London

My Open-Source for Bio-Projects; https://github.com/PawelRosikiewicz

- PvClass Pvthon package for classification & exploration of medical images for Al
- PriceHook Workshop on developing reproducible AI models with real market value
- DataFrame Explorer Dashboard for exploration structured data

SKILLS

Creative, Problem Solving, Team player, Great Organizer

Programming: Python, R, HTML, CSS, SQL, bash, git, Docker

BIG DATA & Machine Learning: scikit-learn, numpy, pandas, Scipy, Tensorflow, Keras,

Data Science: Advanced data analysis and visualizations, Experimental design, A-B testing, Pipeline Development, Dashboards, structured and unstructured data, Statistics, Regression, Classification, data Cleaning, EDA, dealing with missing data, simulated data

Molecular Biology applied to Plant & Fungi: Next Generation Sequencing (Illumina, PacBio, basics of MGI), RNA-seq, RAD-seq, WGS, Amplicon sequencing, Microarrays, microscopy, flow cytometry, FT-IR, qPCR, quantitative Pyrosequencing, microsatellites

Bioinformatics: Variant calling with NGS, fragment length analysis, QC for NGS and SNP data from multiple replicates and libraries, genetic analyses (QC for rare & somatic mutations)

Project management & Product Dev.: Practical experience with developing new well projects, product, MVPs and industrial procedures, Hands-on experience with scaling up R&D protocols to industrial scale. Defined values proposition and product Dev. Roadmap. Helped with business analyses, Market segmentation, Technical sales, and communication.



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Linkedin

www.linkedin.com/in/pawelrosikiewicz/

GitHub

https://github.com/PawelRosikiewicz

My websites

www.SimpleAl.ch www.SwissAl.org

LANGUAGES

Polish, native English, fluent French, intermediate

EDUCATION

Applied Data Science & Machine Learning EPFL-ext 2020

Selected Courses in Machine Learning & Data Science EPFL 2017-2019

Business Development Venturelab, 2018/2019

MSc. Molecular Biology UAM, Poznan, Poland, 2003-2009

CERTIFICATES

Introduction to ML in Production

Practical Introduction to Machine Learning & Deep Learning

Computer skills for biological research

Bayesian Models for Population Genomics

Analysis of differential gene expression

















| pandas











WORK EXPERIENCE Product Developer (Subject Matter Expert) @ MycoSenseAl 2021 Created concept, and product dev. roadmap for Al-powered system helping with the industrial production of edible mushrooms Project Manager @ Dronistics 9.2019 - 6.2021 I worked in a company twice, First, as consultant in 2019, and then, as project manager in 2020-2021 # I led remote R&D team @ Kings College London (see Projects & Leadership section) Provided business Intelligence services to company Helped with product development, technical sales, and marketing. Data Scientist & Business Analyst @ GenomSys SA 11. 2019 - 10.2020 Developed bioinformatic pipelines for analysis of variant calling data from NGS Selected tools and datasets for pipeline development and benchmarking Developed QC steps for new pipeline with Python, GATK, bash, awk, and seqtools Provided direct support to company managers, and board members (R&D & BI projects) Performed feasibility study for new software products, clients and partners Utilized SQL & Python to search large public databases, eg: on genetic tests registry Evaluated market value & requirements for company products, Supported sales & marketing teams Subject Matter Expert in Biotech @ Evologic Technologies GmbH 11.2018 - 4.2020 Implemented, from scratch, new system for industrial production of microorganism's in vitro Conducted several R&D projects for product dev. & management teams Assisted with documentation digitization, and production standards implementation Bioinformatician @ SiB, Swiss Institute of Bioinformatics 12.2015 - 9.2016 Worked on reproducibility problems in BIG DATA from NGS Developed GapJumper; a software used for analysis of genetic markers Bioinformatician @ University of Lausanne 11.2013 - 9.2015 I analyzed variant calling data from 1000+ NGS libraries, (~1TB of raw data) to Studied genetic exchange in polyploid microorganisms ... and genetic& phenotypic stability in microorganisms used as biofertilizers Developed & benchmarked new algorithms for analysis of variant calling data: for parent-offspring analysis in clonal & polyploid organisms detection of somatic mutations Molecular Biologist @ University of Lausanne, 9.2009 - 11.2013 Designed, & Managed several large-scale research projects budget up to 300.000CHF & managed the team of up to 5 people Utilized combination of fluorescent microscopy, in vitro techniques, molecular biology (eg: variant calling, WGS, RNAseq, RAD-seq, qPCR, amplicon sequencing), Bioinformatics and Statistics, to conduct experiments and analyze the data on soil microorganisms MY SELECTED PROJECTS WITH PRACTICAL APPLICATIONS TO INDUSTRY Led the laboratory team for genome sequencing on PacBio platform Publication https://doi.org/10.1073/pnas.1313452110 - cited over 500x! Created new strains of microorganisms that were later used commercially Publication https://doi.org/10.1007/s00572-017-0763-2 Developed cost-efficient protocol for generating RAD-seq markers with illumina HiSeq) The protocol was used to prepare at least 6000 NGS libraries in Switzerland, Israel and Mexico Publication with first use: https://doi.org/10.1038/ismej.2016.29 2.2009 - 9.2009Molecular Biologist @ Institut National de Recherche Agronomique, INRA, Nancy, France

Studied gene expression in microorganisms with custom microarrays (Nimblegene), qPCR and in vitro techniques.

2006 - 2009

Molecular Biologist @ PAN, Polish Academy of Science

- Used RFLP, microsatellites, 18sRNA, microscopy, and in vitro methods to study mycorrhizal fungi diversity in naturally and artificially restored forests, on agral soil.
- Worked on improving of reproducibility of genetic markers

Internships in Molecular & Experiment Biology focused on Fungi in Agronomy

Summer 2008 Summer 2007

- @ UCD Dublin, Ireland, studied gene expression in wheat exposed to toxins from plant pathogens
- @ Lomonosov University, Akhangielsk, Russia, conducted research projects in Russian taiga forest on mycorriza

NGOs

from 2017

Founder & Team Leader @ SwissAl

"Active community of over 4000 AI & Data science professionals"

- I founded and managed all activities of SwissAl for several years:
 - Managed team of up to 8 people & Organized frequent public events, & workshops
 - Responsible for fund raising, speaker recruiting & hiring employees,
 - Marketing for SwissAl, eq. in social media, movie recording, interviews.
 - Facilitated knowledge exchange between academia and industry
 - LINK: www.SwissAl.org